



1646

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/551,341

DATE: 01/24/2002
 TIME: 12:54:53

Input Set : A:\PTO.VSK.txt
 Output Set: N:\CRF3\01242002\I551341.raw

**Does Not Comply
 Corrected Diskette Needed**

SEQUENCE LISTING

```

5 (1) GENERAL INFORMATION:
C--> 7   (i) APPLICANT: Brass, Clifford A., Glue, Paul W., and Piken,
8       Edward.
10   (ii) TITLE OF INVENTION: HCV Combination Therapy
12   (iii) NUMBER OF SEQUENCES: 4
14   (iv) CORRESPONDENCE ADDRESS:
15       (A) ADDRESSEE: Schering-Plough Corporation
16       (B) STREET: 2000 Galloping Hill Road
17       (C) CITY: Kenilworth
18       (D) STATE: New Jersey
19       (E) COUNTRY: USA
20       (F) ZIP: 07033
22   (v) COMPUTER READABLE FORM:
23       (A) MEDIUM TYPE: Floppy disk
24       (B) COMPUTER: Macintosh
25       (C) OPERATING SYSTEM: 7.5.3
C--> 27   (vi) CURRENT APPLICATION DATA:
C--> 28       (A) APPLICATION NUMBER: US/09/551,341
C--> 29       (B) FILING DATE: 18-Apr-2000
30       (C) CLASSIFICATION:
32   (viii) ATTORNEY/AGENT INFORMATION:
33       (A) NAME: Hoffman, Thomas D.
34       (B) REGISTRATION NUMBER: 28,221
35       (C) REFERENCE/DOCKET NUMBER: IN01023K
37   (ix) TELECOMMUNICATION INFORMATION:
38       (A) TELEPHONE: 908-298-5037
39       (B) TELEFAX: 908-298-5388
42 (2) INFORMATION FOR SEQ ID NO: 1:
45   (i) SEQUENCE CHARACTERISTICS:
46       (A) LENGTH: 22 base pairs
47       (B) TYPE: nucleic acid
48       (C) STRANDEDNESS: single
49       (D) TOPOLOGY: linear
W--> 51   (ii) MOLECULE TYPE: DNA (oligonucleotide)
55   (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
59 GTGGTCTGCG GAACCGGTGA GT 22
62 (2) INFORMATION FOR SEQ ID NO: 2:
64   (i) SEQUENCE CHARACTERISTICS:
65       (A) LENGTH: 20 base pairs
66       (B) TYPE: nucleic acid
67       (C) STRANDEDNESS: single
68       (D) TOPOLOGY: linear

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W--> 70 (ii) MOLECULE TYPE: DNA (oligonucleotide)
74 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
76 TGCACGGTCT ACGAGACCTC 20
79 (2) INFORMATION FOR SEQ ID NO: 3:
81 (i) SEQUENCE CHARACTERISTICS:
82 (A) LENGTH: 21 base pairs
83 (B) TYPE: nucleic acid
84 (C) STRANDEDNESS: single
85 (D) TOPOLOGY: linear
W--> 87 (ii) MOLECULE TYPE: DNA (oligonucleotide)
91 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
93 CTGTGAGGAA CTACTGTCTT C 21
97 (2) INFORMATION FOR SEQ ID NO: 4:
99 (i) SEQUENCE CHARACTERISTICS:
100 (A) LENGTH: 21 base pairs
101 (B) TYPE: nucleic acid
102 (C) STRANDEDNESS: single
103 (D) TOPOLOGY: linear
W--> 105 (ii) MOLECULE TYPE: DNA (oligonucleotide)
109 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
111 CCCTATCAGG CAGTACCACA A 21
W--> 115 49
W--> 119 49
W--> 120 21155

VERIFICATION SUMMARY

DATE: 01/24/2002

PATENT APPLICATION: US/09/551,341

TIME: 12:54:54

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L:7 M:220 C: Keyword misspelled or invalid format, [(i) APPLICANT:]
L:27 M:220 C: Keyword misspelled or invalid format, [(vi) CURRENT APPLICATION DATA:]
L:27 M:220 C: Keyword misspelled or invalid format, [(vi) CURRENT APPLICATION DATA:]
L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:51 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:70 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:87 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:105 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4
L:115 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
L:119 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4
L:120 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:4